

Figure 1

1 GATTCGGCAGAGAAAACCTTTTAAATCTTTAGTTATTTCTTAATACTTAGAACACTTAAAC 60

61 AAAACTTTACAAAACAAAAGAGCAGAATAATTAGATCCTTTCAGGAGAATATGACTTTTT 120

121 TTTCTTAAGCACACTGGACCATAGAGGAAGACCAAAGGAATGTACAGTTGCCTGCTCCTT 180  
1 M Y S C L L L 7

181 CCTGACTTGCTGTATTTgACTCTGtCCCCACTGGTGGTGGCAATGCTATTAAACCCACAC 240  
8 P D L L Y L T L S P L V V A M L L T P H 27

241 TTTAACGTGGCAAATCCCCAGAATCTGTTGGCTGGTCTCTGGCTAGAGAATGAGCACAGT 300  
28 F N V A N P Q N L L A G L W L E N E H S 47

301 TTCACCCCTTATGGCTCCAGAAAGAGCAAGAACACACCACTGCCAGCCAGAAGAGAGAAAA 360  
48 F T L M A P E R A R T H H C Q P E E R K 67

361 GTCTTGTTCTGTCTCTTTCCATTGTCCCAATAGCCAAGCACAGGTTCAACCACCCCAA 420  
68 V L F C L F P I V P N S Q A Q V Q P P Q 87

421 ATGCCACCCCTTCTGCTGTGCAGCAGCCAAGGAAAAGACCCAGGAGGAGCAGCTCCAAGAA 480  
88 M P P F C C A A A K E K T Q E E Q L Q E 107

481 CCTCTGGGCAGTCAGTGCCCAGATACTTGCCCCAATTCTTTGTGTCCAAGCCACACTCAG 540  
108 P L G S Q C P D T C P N S L C P S H T Q 127

541 CTGACAAAAGCCAACACTTTGTCTCTCTTTTTTTTTTTTTCTTTTTTTTGGAGCAGAGTT 600  
128 L T K A N T L S L F F F F S F F L S R V 147

601 TCACTCTTGTCACCCAGGCTGGAGTGCAATGgCAGGATCTTGGCTCATTGCAACCTCCAC 660  
148 S L L S P R L E C N G R I L A H C N L H 167

661 CtCCCGGGTTCAAGCAATTCTCCTGTCTCAGCCTCTCGA 699  
168 L P G S S N S P V S A S R 180

Figure 2

